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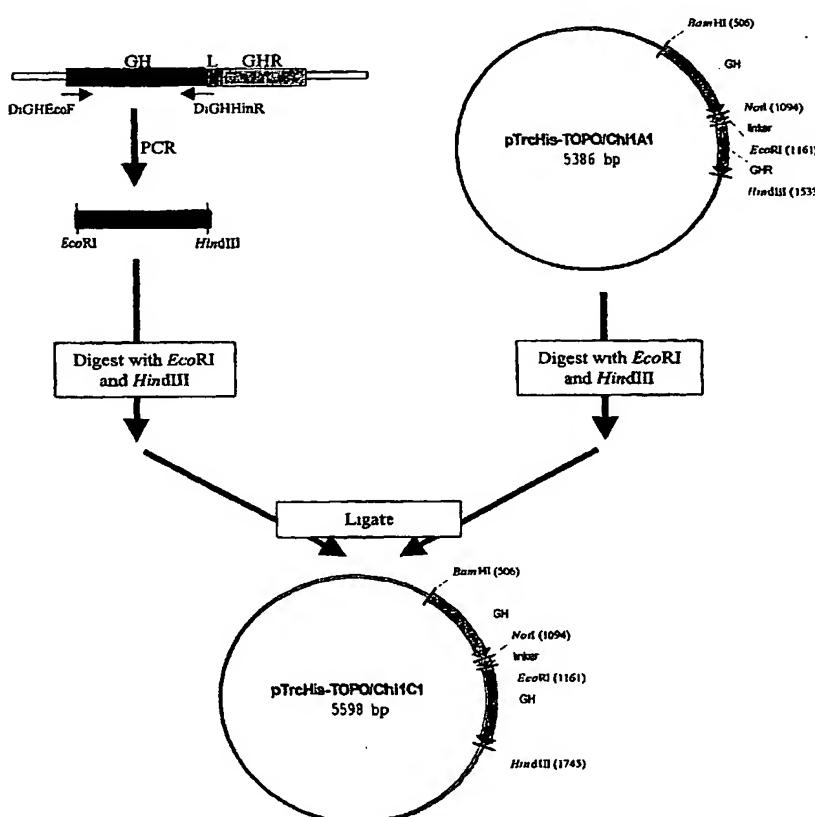
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(54) Title: POLYPEPTIDE VARIANTS



(57) Abstract: The invention relates to the provision of oligomeric polypeptides (dimers, trimers, etc) comprising the ligand binding domains of cytokines which are linked via flexible polypeptide linker molecules. The linker molecules optionally comprise protease sensitive sites to modulate the release of biologically active cytokines when administered to a human or animal subject.

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POLYPEPTIDE VARIANTS

The invention relates to polypeptides which comprise more than two ligand binding domains of a cytokine wherein the domains are linked by a flexible linker which optionally comprises
5 a proteolytic cleavage site.

Ligands which interact with receptors to bring about a suitable biochemical response are known as agonists and those that prevent, or hinder, a biochemical response are known as antagonists. For example, and not by way of limitation, cell specific growth factors are
10 ligands that act as agonists and bind receptors located at the cell surface. Activation of the receptors by ligand-specific binding promotes cell proliferation via activation of intracellular signalling cascades that result in the expression of, amongst other things, cell-cycle specific genes, and the activation of quiescent cells to proliferate. Growth factors may also activate cellular differentiation.

15

A large group of growth factors, referred to as cytokines, are involved in a number of diverse cellular functions. These include, by example and not by way of limitation, modulation of the immune system, regulation of energy metabolism and control of growth and development. Cytokines which are secreted by lymphocytes are termed lymphokines (also known as
20 interleukins). Those secreted by monocytes and macrophages are termed monokines. Cytokines are also secreted by endocrine glands, (for example growth hormone (GH) by the pituitary gland), and adipose cells (for example leptin). Cytokines mediate their effects via receptors expressed at the cell surface of target cells.

25

Receptors of the cytokine receptor family possess a single transmembrane domain and lack intrinsic enzyme activity (Kishimoto *et al.*, 1994). Upon binding of a cytokine to a cognate receptor, either receptor homo- or hetero-dimerisation or oligomerisation occurs. The receptor complex is internalised and signalling occurs through the activation of associated signalling cascades that include the Jak/Stat and Mapk pathways. Internalisation is followed
30 by a recycling step whereby the receptor molecule is regenerated for further use within the cell.

An example of the above is described with respect to GH and its binding to the GHR. This example is merely meant to be illustrative and not limiting and is an example of a cytokine

which activates a signal transduction cascade by binding, dimerisation and internalisation of the receptor:ligand complex. It is known that a single molecule of growth hormone (GH) associates with two identical receptor molecules (Cunningham *et al.*, 1991; de Vos *et al.*, 1992; Sundstrom *et al.*, 1996; Clackson *et al.*, 1998). This occurs through two unique
5 receptor-binding sites on GH and a common binding pocket on the extracellular domain of two receptors. Site 1 on the GH molecule has a higher affinity than site 2, and receptor dimerization is thought to occur sequentially with one receptor binding to site 1 on GH followed by recruitment of a second receptor to site 2.

10 The extracellular portion of the GHR exists as two linked domains each of approximately 100 amino acids (SD-100), the C-terminal SD-100 domain being closest to the cell surface and the N-terminal SD-100 domain being furthest away. It is a conformational change in these two domains that occurs on hormone binding with the formation of the hetero-trimeric complex GHR-GH-GHR. It has been proposed that ligand-driven receptor dimerization is the key event
15 leading to signal activation (Cunningham *et al.*, 1991), triggering phosphorylation cascades that include the Jak2/Stat5 pathway (Argetsinger & Carter-Su, 1996). Using confocal microscopy and Fluorescence Resonance Energy Transfer (FRET) it is known that there is very rapid internalisation of GHR after ligand binding and that internalisation and signaling are independent functions (Maamra *et al.*, 1999). Internalisation of the GHR-GH-GHR
20 complex is followed by a recycling step whereby the receptor molecule is regenerated for further use within the cell.

Examples of cytokines which are related to GH are leptin and erythropoietin (EPO). The leptin receptor and the EPO receptor share considerable structural homology with the GHR
25 and require a similar dimerisation process to trigger signalling. Leptin suppresses appetite and leptin resistance is associated with obesity. A leptin receptor antagonist will provide a treatment for anorexia nervosa. EPO excess causes polycythaemia which may be secondary to hypoxia (chronic lung disease), or primary in the case of polycythaemia rubra vera (a disorder of excess red blood cells). An EPO antagonist will provide a therapy for polycythaemia.

30

The disorders of acromegaly and gigantism result from an excess of growth hormone, usually due to pituitary tumours. A drug currently under trial is the pegylated GH antagonist B2036, designed using recently acquired knowledge of the molecular structure of the GHR.

Unfortunately, to antagonise GH action very high levels of B2036 are required, over a 1000 times higher than endogenous GH levels.

5 The invention relates, *inter alia*, to the provision of oligomeric polypeptides (dimers, trimers etc) comprising the ligand binding domains of cytokines which are linked via flexible polypeptide linker molecules which optionally include protease sensitive sites to modulate the release of biologically active cytokines when administered to an animal.

10 According to a first aspect of the invention there is provided a polypeptide comprising more than two ligand binding domains of a cytokine receptor wherein said domains are linked by a linker molecule.

Preferably the linker molecule comprises at least one proteolytic cleavage site.

15 Preferably said cleavage site is sensitive to a serum protease.

Preferably said cleavage site comprises the amino acid sequence: LVPRGS, or variant thereof.

20 In a further preferred embodiment of the invention said cleavage site comprises at least one copy of the amino acid sequence: SGGGG, or functional variant thereof. Preferably, said cleavage site comprises the amino acid sequence PGISGGGGGG.

25 More preferably still said cleavage site comprises the amino acid sequence: LVPRGS PGISGGGGGG, or variant thereof.

Alternatively, said cleavage site comprises at least two copies of the amino acid sequence SGGGG, or functional variant thereof, which flank said cleavage site.

30 In a further preferred embodiment of the invention said cleavage site is sensitive to the serum protease thrombin.

In a further preferred embodiment of the invention said polypeptide comprises a plurality of ligand binding domains. Preferably said polypeptide has 3, 4, 5, 6, 7, 8, 9,

or 10 ligand binding domains. Preferably said polypeptide has greater than 10 ligand binding domains.

5 In a further preferred embodiment of the invention said polypeptide has 4, 6, 8, 10, or 12 ligand binding domains.

In a preferred embodiment of the invention said polypeptide comprises at least four ligand binding domains.

10 In a further embodiment of the invention said polypeptide is an antagonist.

In an alternative preferred embodiment of the invention said polypeptide is an agonist.

15 In a further preferred embodiment of the invention said ligand binding domain is selected from the ligand binding domains of the cytokines selected from the group consisting of: growth hormone; leptin; erythropoietin; prolactin; interleukins (IL), IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-9, IL-10, IL-11; the p35 subunit of IL-12, IL-13, IL-15; granulocyte colony stimulating factor (G-CSF); granulocyte macrophage colony stimulating factor (GM-CSF); ciliary neurotrophic factor (CNTF); cardiotrophin-1
20 (CT-1); leukaemia inhibitory factor (LIF); oncostatin M (OSM); interferon, IFN α and IFN γ .

In a preferred embodiment of the invention said ligand binding domain is the ligand binding domain of growth hormone.

25 A single hGH molecule binds to two GHR molecules. The hGH molecule interacts with one receptor molecule through a high affinity site, and with the other through a low affinity site. A single protein chain consisting of hGH linked to hGH will contain two high affinity sites which can bind strongly to a pair of receptor molecules and two
30 low affinity sites.

In one embodiment of the invention, two or more copies of a ligand binding domain are expressed on a single polypeptide chain and the sequence of the tandem or

oligomeric cytokine is arranged ‘ligand binding domain-linker-ligand binding domain’.

5 In a further preferred embodiment of the invention said ligand binding domain is the binding domain of leptin.

Preferably the linker comprises at least one copy of the peptide:

Gly Gly Gly Gly Ser Ser Ser Ser (hereinafter referred to as “Gly4Ser4”).

10 In one embodiment of the invention the linker is 8 amino acids in length and consists of one copy of the Gly4Ser4 linker. In an alternative embodiment of the invention, the linker is 16 amino acids in length and consists of two copies of the Gly4Ser4 linker. In a further alternative embodiment of the invention said linker is 24 amino
15 acids in length and consists of three copies of the Gly4Ser4 linker.

More preferably the linker is a polypeptide which comprises 5 to 50 amino acid residues. Most preferably the linker comprises 5 to 30 amino acid residues.

20 Most preferably the linker comprises at least one copy of the peptide:

Gly Gly Gly Gly Ser (hereinafter referred to as “(Gly4Ser)” or (G₄S)).

25 In one embodiment of the invention the linker is 5 amino acids in length and consists of one copy of the (Gly4Ser) linker. In an alternative embodiment of the invention, the linker is 10 amino acids in length and consists of two copies of the (Gly4Ser)₂ linker. In a further alternative embodiment of the invention said linker is 15 amino acids in length and consists of three copies of the (Gly4Ser) linker. In a further alternative embodiment of the invention said linker is 20 amino acids in length and
30 consists of 4 copies of the (Gly4Ser)₄ linker.

In an alternative embodiment of the invention, the polypeptide is a fusion protein comprising inframe translational fusions of ligand binding domains according to the invention.

It will be apparent to one skilled in the art that alternative linkers can be used to link ligand binding domains, for example the use of chemical protein crosslinkers. For example homo-bifunctional crosslinker such as disuccinimidyl-suberimidate-dihydrochloride; dimethyl-adipimide-dihydrochloride; 1,5,-2,4 dinitrobenezene or 5 hetero-bifunctional crosslinkers such as N-hydroxysuccinimidyl 2, 3-dibromopropionate; 1-ethyl-3-[3-dimethylaminopropyl] carbodiimide hydrochloride; succinimidyl 4-[n-maleimidomethyl]-cyclohexane-1-carboxylate.

Further examples of chemical crosslinks include the provision of chemically modified 10 linker molecules and/or ligand binding domains. For example, if one end of the linker molecule is terminated with an amino terminal lysine residue and the ligand binding with a carboxyl-terminal glutamine residue then ligand binding domains can be oligomerised with transglutaminase.

15 According to a further aspect of the invention there is provided a nucleic acid molecule comprising a nucleic acid sequence which encodes a polypeptide according to the invention.

In a preferred embodiment of the invention said nucleic acid molecule comprises a nucleic acid sequence selected from the group consisting of:

20 i) a sequence represented by Figs 4 or 6;

ii) a sequence which hybridises to the sequence of (i) above and which has cytokine receptor modulating activity; and

iii) a sequence which is degenerate as a result of the genetic code to the sequences defined in (i) and (ii) above.

25 In a preferred embodiment of the invention said nucleic acid hybridises under stringent hybridisation conditions to the sequences represented in Figs 4 or 6.

It is well known in the art that optimal hybridisation conditions can be calculated if the sequence of the nucleic acid is known. For example, hybridisation conditions can be determined by the GC content of the nucleic acid subject to hybridisation. Please

see Sambrook *et al.* (1989) Molecular Cloning; A Laboratory Approach. A common formula for calculating the stringency conditions required to achieve hybridisation between nucleic acid molecules of a specified homology is:

5 $T_m = 81.5^0 C + 16.6 \log [Na^+] + 0.41[\% G + C] - 0.63 (\%formamide).$

Typically, hybridisation conditions uses 4 – 6 x SSPE (20x SSPE contains 175.3g NaCl, 88.2g NaH₂PO₄ H₂O and 7.4g EDTA dissolved to 1 litre and the pH adjusted to 7.4); 5-10x Denhardts solution (50x Denhardts solution contains 5g Ficoll (type 400,

10 Pharmacia), 5g polyvinylpyrrolidone abd 5g bovine serum albumen/500ml; 100μg-1.0mg/ml sonicated salmon/herring DNA; 0.1-1.0% sodium dodecyl sulphate; optionally 40-60% deionised formamide. Hybridisation temperature will vary depending on the GC content of the nucleic acid target sequence but will typically be between 42⁰- 65⁰ C.

15 According to a further aspect of the invention there is provided a polypeptide which is encoded by a nucleic acid molecule according to the invention.

In a preferred embodiment of the invention the polypeptide so encoded is modified by deletion, addition or substitution of at least one amino acid residue. Ideally said modification enhances the antagonistic or agonistic effects of said polypeptide with
20 respect to the inhibition or activation of receptor mediated cell signalling.

Alternatively, or preferably, said modification includes the use of modified amino acids in the production of recombinant or synthetic forms of polypeptides.

It will be apparent to one skilled in the art that modified amino acids include, by way of example and not by way of limitation, 4-hydroxyproline, 5-hydroxylysine, N⁶-acetyllysine, N⁶-methyllysine, N⁶,N⁶-dimethyllysine, N⁶,N⁶,N⁶-trimethyllysine, cyclohexyalanine, D-amino acids, ornithine. The incorporation of modified amino acids may confer advantageous properties on polypeptides. For example, the incorporation of modified amino acids may increase the affinity of the polypeptide for its binding site, or the modified amino acids may confer increased *in vivo* stability on
30 the polypeptide thus allowing a decrease in the effective amount of therapeutic polypeptide administered to a patient.

According to a yet further aspect of the invention there is provided a vector including a DNA molecule according to any preceding aspect or embodiment of the invention.

5 In a preferred embodiment of the invention said vector is provided with means to manufacture recombinantly the polypeptide of the invention.

In a preferred embodiment of the invention said vector is an expression vector adapted for prokaryotic gene expression.

10 Prokaryotic expression systems are well known in the art and comprise vectors adapted for high level constitutive and inducible expression. Inducible expression systems are particularly advantageous if the recombinant polypeptide is toxic to the bacterial cell. Induction of expression is tightly regulated by promoters responsive to various inducers (e.g. IPTG inducible). Bacterial cells can be grown to stationary 15 phase before induction thereby reducing harmful effects of toxic polypeptides.

20 Additionally it is also well known in the art that certain polypeptides are difficult to manufacture recombinantly due, for example, to protein instability or problems of aggregation. It is well known that genetically modified bacterial strains are available which are mutated in genes (e.g. bacterial proteases) which facilitate the production of native and recombinant bacterial polypeptides.

25 In a further preferred embodiment of the invention said vector is an expression vector adapted for eukaryotic gene expression.

Typically said adaptation includes, by example and not by way of limitation, the provision of transcription control sequences (promoter sequences) which mediate cell/tissue specific expression. These promoter sequences may be cell/tissue specific, inducible or constitutive.

30 Promoter is an art recognised term and, for the sake of clarity, includes the following features which are provided by example only, and not by way of limitation. Enhancer elements are *cis* acting nucleic acid sequences often found 5' to the transcription initiation site of a gene (enhancers can also be found 3' to a gene sequence or even

located in intronic sequences and are therefore position independent). Enhancers function to increase the rate of transcription of the gene to which the enhancer is linked. Enhancer activity is responsive to *trans* acting transcription factors (polypeptides) which have been shown to bind specifically to enhancer elements. The 5 binding/activity of transcription factors (please see Eukaryotic Transcription Factors, by David S. Latchman, Academic Press Ltd., San Diego) is responsive to a number of environmental cues which include, by example and not by way of limitation, intermediary metabolites (eg glucose, lipids), environmental effectors (eg light, heat).

10 Promoter elements also include so called TATA box and RNA polymerase initiation selection (RIS) sequences which function to select a site of transcription initiation. These sequences also bind polypeptides which function, *inter alia*, to facilitate transcription initiation selection by RNA polymerase.

15 Adaptations also include the provision of selectable markers and autonomous replication sequences which both facilitate the maintenance of said vector in either the eukaryotic cell or prokaryotic host. Vectors which are maintained autonomously are referred to as episomal vectors.

20 Adaptations which facilitate the expression of vector encoded genes include the provision of transcription termination/polyadenylation sequences. This also includes the provision of internal ribosome entry sites (IRES) which function to maximise expression of vector encoded genes arranged in bicistronic or multi-cistronic expression cassettes.

25 These adaptations are well known in the art. There is a significant amount of published literature with respect to expression vector construction and recombinant DNA techniques in general. Please see, Sambrook *et al.* (1989) Molecular Cloning: A Laboratory Manual, Cold Spring Harbour Laboratory, Cold Spring Harbour, NY and references therein; Marston, F (1987) DNA Cloning Techniques: A Practical Approach Vol. III IRL Press, Oxford UK; DNA Cloning: F M Ausubel *et al.*, Current Protocols in Molecular Biology, John Wiley & Sons, Inc.(1994).

In yet a further aspect of the invention there is provided a method to prepare a polypeptide according to the invention comprising:

- 5 (i) growing a cell transformed or transfected with a nucleic acid or vector of the present invention in conditions conducive to the manufacture of said polypeptide; and
- (ii) purifying said polypeptide from said cell, or its growth environment.

10 In a preferred method of the invention said vector encodes, and thus said recombinant polypeptide is provided with, a secretion signal to facilitate purification of said binding agent polypeptide.

In yet a further aspect of the invention there is provided a cell transformed/transfected with the vector or nucleic acid according to the invention.

15

Preferably said cell eukaryotic and is selected from: fungal; insect (e.g. *Spodoptera frugiperda*); amphibian; plant; mammalian.

20

According to a further aspect of the invention there is provided the use of the polypeptide according to the invention as a pharmaceutical. Preferably there is provided a pharmaceutical composition comprising the polypeptide according to the invention. Preferably said pharmaceutical composition includes a carrier, excipient and/or a diluent.

25 In a further preferred embodiment of the present invention said polypeptide is used for the manufacture of a medicament for use in the treatment of a disease selected from the group consisting of: acromegaly; gigantism; GH deficiency; Turners syndrome; renal failure; osteoporosis; diabetes mellitus; cancer; obesity; insulin resistance; hyperlipidaemia; hypertension; anaemia; autoimmune and infectious disease; inflammatory disorders including rheumatoid arthritis.

The invention also provides for a method of treating a human or animal subject comprising administering an effective amount of the polypeptide, pharmaceutical composition or medicament to said subject.

5 The polypeptides or compositions of the invention can be administered by any conventional route, including injection or by gradual infusion over time. The administration may, for example, be oral, intravenous, intraperitoneal, intramuscular, intracavity, subcutaneous, transdermal or delivered by a non-pathological GMO engineered to secrete the polypeptide.

10

The pharmaceutical compositions used in the foregoing methods preferably are sterile and contain an effective amount of the polypeptide according to the invention for producing the desired response in a unit of weight or volume suitable for administration to a patient.

15

When administered, the pharmaceutical preparations of the invention are applied in pharmaceutically-acceptable amounts and in pharmaceutically-acceptable compositions. The term "pharmaceutically acceptable" means a non-toxic material that does not interfere with the effectiveness of the biological activity of the active 20 ingredients. Such preparations may routinely contain salts, buffering agents (e.g., acetic acid in a salt; citric acid in a salt; boric acid in a salt; and phosphoric acid in a salt), preservatives (e.g., benzalkonium chloride; chlorobutanol; parabens and thimerosal, compatible carriers, and optionally other therapeutic agents.

20

25 Compositions may be combined, if desired, with a pharmaceutically-acceptable carrier. The term "pharmaceutically-acceptable carrier" as used herein means one or more compatible solid or liquid fillers, diluents or encapsulating substances which are suitable for administration into a human. The term "carrier" denotes an organic or inorganic ingredient, natural or synthetic, with which the active ingredient is 30 combined to facilitate the application.

The pharmaceutical compositions may conveniently be presented in unit dosage form and may be prepared by any of the methods well-known in the art of pharmacy. All methods include the step of bringing the active agent into association with a carrier

which constitutes one or more accessory ingredients. In general, the compositions are prepared by uniformly and intimately bringing the active compound into association with a liquid carrier, a finely divided solid carrier, or both, and then, if necessary, shaping the product.

5

Compositions suitable for oral administration may be presented as discrete units, such as capsules, tablets, lozenges, each containing a predetermined amount of the active compound. Other compositions include suspensions in aqueous liquids or non-aqueous liquids such as a syrup, elixir or an emulsion.

10

Compositions suitable for parenteral administration conveniently comprise a sterile aqueous or non-aqueous preparation of polypeptides which is preferably isotonic with the blood of the recipient. This preparation may be formulated according to known methods using suitable dispersing or wetting agents and suspending agents. The

15

sterile injectable preparation also may be a sterile injectable solution or suspension in a non-toxic parenterally-acceptable diluent or solvent, for example, as a solution in 1,3-butane diol. Among the acceptable vehicles and solvents that may be employed are water, Ringer's solution, and isotonic sodium chloride solution. In addition, sterile, fixed oils are conventionally employed as a solvent or suspending medium.

20

For this purpose any bland fixed oil may be employed including synthetic mono-or di-glycerides. In addition, fatty acids such as oleic acid may be used in the preparation of injectables. Carrier formulation suitable for oral, subcutaneous, intravenous, intramuscular, etc. administrations can be found in Remington's Pharmaceutical Sciences, Mack Publishing Co., Easton, PA.

25

According to a further aspect of the invention there is provide a method of treatment comprising administering to an animal, preferably a human, an effective amount of the nucleic acid or vector according to the invention.

30

An embodiment of the invention will now be described by example only and with reference to the following figures wherein;

Figure 1 illustrates a plasmid map of pTrcHis χ 1A1.

Figure 2 illustrates primers used in the synthesis of the tandem constructs.

Figure 3 is a schematic diagram for the construction of pTrcHis χ 1C1. pTrcHis χ 1C2

5 is constructed in the same way, however the forward primer used is DiGHNotF and the restriction enzymes used are *NotI* and *HindIII*.

Figure 4 illustrates the DNA sequence for growth hormone tandem segregated by a

thrombin cleavable linker. The linker region is shown in italics, with the thrombin

10 cleavage site in bold.

Figure 5 illustrates the protein sequence for growth hormone tandem segregated by a

thrombin cleavable linker. The linker region is shown in italics, with the thrombin

cleavage site in bold.

15

Figure 6 illustrates the DNA sequence for leptin tandem segregated by a thrombin cleavable linker. The linker region is shown in italics, with the thrombin cleavage site in bold.

20

Figure 7 illustrates the protein sequence for leptin tandem segregated by a thrombin cleavable linker. The linker region is shown in italics, with the thrombin cleavage site in bold. The cleavable linker is optional in this construct.

25

Figure 8 shows the Coomassie stained SDS-PAGE gel and western blot of the purified GH-tandem (χ 1C1).

Figure 9 is the bioassay data generated for χ 1C1, with data for in-house synthesised

GH and commercially produced GH for comparison. The activity of the tandem is

similar to the in-house generated GH. In each case the concentrations of the test

30 protein used were 0ng, 6.25ng, 12.5ng, 25ng, 50ng and 100ng.

Figure 10 is a schematic diagram illustrating the construction of pTrcHis χ 2C1.

Figure 11 shows the western blot of the SDS-PAGE gel on which the proteins expressed by clones of *E. coli* SURE:pTrcHis χ 2C1 cells were run. The expected size of the leptin-tandem is ~37kDa and a faint band is observed at this size. The major band however is approximately half of this size. This suggests that the leptin tandem is being expressed but is most probably cleaved to the single leptin domains.

Materials and Methods

The sequence encoding GHR in pTrcHis χ 1A1 (Fig. 1) was replaced with another GH gene (residues 1-191) to produce two successive GH genes linked with a (G₄S)₄ linker. The resultant construct (pTrcHis χ 1C1) was transformed into *E. coli* SURE cells, a DNA recombination deficient strain of *E. coli*. Clones expressing GH-(G₄S)₄-GH protein were identified by western blotting using anti-GH (10A7, mouse IgG1) probed with Sheep anti-mouse-HRP (Amersham). Another GH-tandem, which lacked the (G₄S)₄ linker (GH-GH) was also constructed (pTrcHis χ 1C2) using the same method.

GH-tandem protein was purified from cell lysates using a metal chelate affinity column (Probond resin, Invitrogen) followed by an ion exchange column (MonoQ, Pharmacia).

The effect of the GH-tandem proteins were analysed using an established bioassay (Ross *et al.*, 1997).

The leptin gene was originally cloned into pHEAT; a temperature-inducible vector. However for expression in *E. coli* SURE cells the gene was sub-cloned into the pTrcHis plasmid. A (G₄S)₄ linker was then introduced and finally the second leptin domain was ligated into the gene to produce the construct that would express the leptin tandem. This construct pTrcHis χ 2C1 was then transformed into *E. coli* SURE cells. Expression of the leptin-tandem was verified by western blot using anti-leptin antibodies (Sigma), developed in rabbit, probed with anti-rabbit-HRP (Sigma).

Cloning of the GH-Tandems

PCR was used to produce a fragment of DNA which consisted of a restriction site (*NotI* or *EcoRI*), followed by the GH gene and then a *HindIII* restriction site. This was ligated into pTrcHis χ 1A1, which had been digested with the relevant restriction enzymes, to produce the constructs χ 1C1 (GH-(G₄S)₄-GH) and χ 1C2 (GH-GH). The primers used for the PCR's are shown in Fig 2, and the reaction scheme is shown in Fig 3.

Purification of the GH-Tandems

10

Induced cells (resuspended in 20mM sodium phosphate buffer, 500mM sodium chloride, pH 7.8) were lysed with a combination of 100 μ g/ml (final concentration) hen egg white lysozyme and sonication. Insoluble material was removed by centrifugation at 4000rpm for 20 minutes.

The cleared cell lysate was applied to a 5ml Probond resin column (Invitrogen), equilibrated with 20mM sodium phosphate buffer, 500mM sodium chloride, 5% glycerol, pH 7.8. The column was then washed with 10 column volumes of 20mM sodium phosphate buffer, 500mM sodium chloride, 5% glycerol, pH 6.0. Bound protein was eluted using 5ml 20mM sodium phosphate buffer, 500mM sodium chloride, 5% glycerol, 500mM imidazole, pH 6.0.

20 The protein was dialysed overnight against Low Salt Buffer (25mM TRIS, 1mM EDTA, 5% glycerol, pH 8.0) and then centrifuged to remove any particulate matter.

25 The protein sample was then loaded onto a Mono-Q column (Pharmacia), which had been pre-equilibrated with Low Salt Buffer. After a 10 column volume wash with Low Salt Buffer, the bound proteins were eluted over 20 column volumes using a gradient between 0M sodium chloride to 1M sodium chloride (in 25mM TRIS, 1mM EDTA, 5% glycerol, pH8.0). Peaks on the elution profile were analysed by SDS-PAGE and western blotting.

30 GH-tandem protein was then concentrated (if required) using a Amicon Centriprep Y-10 column.

The purity of the purified χ 1C1 was confirmed by SDS-PAGE, by both coomassie staining and western blot (Fig. 8). Once the integrity of this sample had been confirmed, χ 1C1 was submitted to the previously established bioassay (Ross *et al.*, 5 1997) (Fig. 9).

Cloning of the Leptin-Tandems

PCR using the primers Lep2TrcFOR and Lep2TrcREV (Fig 2) was used to generate 10 DNA sequence consisting of (*Nhe*I)-LEPTIN-(*Not*I)-(X*ho*I)-(S*a*II)-STOP-(*Eco*RI) from pHEATLeptin. The terminal restriction sites were introduced by PCR and the internal restriction sites were already present in the pHEATLeptin vector. The PCR product generated was ligated into pTrcHis χ 1A1 between *Nhe*I and *Eco*RI restriction sites to produce pTrcHisLeptin.

15

PCR was then used to generate the (G₄S)₄-encoding linker flanked by *Not*I and X*ho*I restriction sites, the primers used were LepLinkFOR and LepLinkREV (Fig 2). This was ligated into pTrcHisLeptin between the *Not*I and X*ho*I sites to produce pTrcHisLepLink.

20

The second leptin gene flanked by X*ho*I and S*a*II restriction sites was generated by PCR, using the primers Lep2FOR and Lep2REV (Fig 2). This was ligated between the X*ho*I and S*a*II restriction sites to produce the construct which would express the leptin tandem, pTrcHis χ 2C1. This process for the generation of the leptin-tandem is 25 shown in Fig. 10.

This plasmid was transformed into *E. coli* SURE cells and expression studies carried out, visualisation of expression was performed by western blot (Fig. 11)

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25

30

CLAIMS

1. A polypeptide comprising more than two ligand binding domains of a cytokine receptor wherein said domains are linked by a linker molecule and wherein the linker molecule comprises at least one proteolytic cleavage site.
5
2. A polypeptide according to claim 1 wherein said cleavage site is sensitive to a serum protease.
- 10 3. A polypeptide according to claim 2 wherein the serum protease is thrombin.
4. A polypeptide according to any of the preceding claims wherein said cleavage site comprises the amino acid sequence LVPRGS, or a variant thereof.
- 15 5. A polypeptide according to any of the preceding claims wherein said cleavage site comprises the amino acid sequence SGGGG, or a variant thereof.
6. A polypeptide according to any of the preceding claims wherein said cleavage site comprises the amino acid sequence PGISGGGGGG.
- 20 7. A polypeptide according to any of the preceding claims wherein said cleavage site comprises the amino acid sequence: LVPRGSPGISGGGGGG, or a variant thereof.
- 25 8. A polypeptide according to any of the preceding claims wherein said cleavage site comprises at least two copies of the amino acid sequence SGGGG, or a variant thereof, which flank said cleavage site.
9. A polypeptide according to any of the preceding claims wherein said polypeptide comprises at least four ligand binding domains.
30
10. A polypeptide according to claim 9 wherein said polypeptide has 4, 6, 8, 10, or 12 ligand binding domains.

11. A polypeptide according to any of the preceding claims 1 to 8 wherein said polypeptide has 3, 4, 5, 6, 7, 8, 9, or 10 ligand binding domains.

12. A polypeptide according to any of the preceding claims 1 to 9 wherein said 5 polypeptide has greater than 10 ligand binding domains.

13. A polypeptide according to any of the preceding claims wherein said polypeptide is an antagonist.

10 14. A polypeptide according to any of the preceding claims wherein said polypeptide is an agonist.

15 15. A polypeptide according to any of the preceding claims wherein said cytokine receptor ligand binding domain is selected from the ligand binding domains of the cytokines selected from the group consisting of: growth hormone; leptin; erythropoietin; prolactin; interleukins (IL), IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-9, IL-10, IL-11; the p35 subunit of IL-12, IL-13, IL-15; granulocyte colony stimulating factor (G-CSF); granulocyte macrophage colony stimulating factor (GM-CSF); ciliary neurotrophic factor (CNTF); cardiotrophin-1 (CT-1); leukaemia inhibitory factor (LIF); oncostatin M (OSM); interferon, IFN α and IFN γ .
20

16. A polypeptide according to claim 15 wherein the binding domain is the ligand binding domain of growth hormone.

25 17. A polypeptide according to claim 16 wherein the binding domain is the ligand binding domain of leptin.

18. A polypeptide according to any of the preceding claims wherein the linker is a polypeptide which comprises from 5 to 50 amino acid residues.

30 19. A polypeptide according to claim 18 wherein the linker comprises from 5 to 30 amino acid residues.

20. A polypeptide according to any of the preceding claims wherein the linker comprises at least one copy of the peptide Gly Gly Gly Gly Ser.

21. A polypeptide according to claim 20 wherein the linker is 5 amino acids in
5 length and consists of one copy of the (Gly4Ser) linker.

22. A polypeptide according to claim 20 wherein the linker is 10 amino acids in length and consists of two copies of the (Gly4Ser)2 linker.

10 23. A polypeptide according to claim 20 wherein the linker is 15 amino acids in length and consists of three copies of the (Gly4Ser) linker.

24. A polypeptide according to claim 20 wherein the linker is 20 amino acids in length and consists of 4 copies of the (Gly4Ser)4 linker.

15 25. A polypeptide according to any of the preceding claims wherein the polypeptide is a fusion protein comprising inframe translational fusions of ligand binding domains.

20 26. A polypeptide according to any of the preceding claims comprising chemical crosslinkers wherein the chemical crosslinkers serve to link the ligand binding domains.

25 27. A polypeptide according to claim 26 wherein the chemical crosslinker comprises a homo-bifunctional crosslinker selected from the group consisting of; disuccinimidyl-suberimidate-dihydrochloride; dimethyl-adipimidate-dihydrochloride; 1,5,-2,4 dinitrobenezene.

30 28. A polypeptide according to claim 26 or claim 27 wherein the crosslinker comprises a hetero-bifunctional crosslinker selected from the group consisting of; N-hydroxysuccinimidyl 2, 3-dibromopropionate; 1-ethyl-3-[3-dimethylaminopropyl] carbodiimide hydrochloride; succinimidyl 4-[n-maleimidomethyl]-cyclohexane-1-carboxylate.

29. A nucleic acid molecule comprising a nucleic acid sequence which encodes a polypeptide according to any of the preceding claims.

30. A nucleic acid molecule comprising the sequence selected from the group consisting of:

5 (i) the sequence represented by Figs 4 or 6;

(iii)a sequence which hybridises to the sequence of (i) above and which has cytokine receptor modulating activity; and

(iii)a sequence which is degenerate as a result of the genetic code to the sequences defined in (i) and (ii) above.

10 31. A nucleic acid molecule which hybridises under stringent hybridisation conditions to the sequences represented in Figs 4 or 6.

32. A polypeptide encoded by the nucleic acid molecule according to any of claims 29 to 31.

15 33. A polypeptide according to claim 32 wherein said polypeptide is modified by deletion, addition, and/or substitution of at least one amino acid residue and said modification enhances the antagonistic or agonistic effects of said polypeptide with respect to the inhibition or activation of receptor mediated cell signalling.

34. A vector including a DNA molecule wherein said DNA molecule encodes a polypeptide according to any of claims 1 to 28, 32 or 33.

20 35. A vector according to claim 34 wherein said vector is an expression vector adapted for prokaryotic or eukaryotic gene expression.

25 36. A vector according to claim 34 or 35 wherein said vector encodes, and thus said recombinant polypeptide is provided with, a secretion signal to facilitate purification of said polypeptide.

37. A method to prepare a polypeptide according to any of claims 1 to 28, 32 or 33 comprising;

- (i) growing a cell transformed or transfected with a nucleic acid of any of claims 29 to 31 or a vector of claims 34 to 36 in conditions conducive to the manufacture of said polypeptide; and
- 5 (ii) purifying said polypeptide from said cell, or its growth environment.

38. A cell transformed/transfected with the vector of claims 34 to 36 or the nucleic acid of claims 29 to 31.

10 39. Use of the polypeptide of any of claims 1 to 28, 32 or 33 and/or the nucleic acid of claims 29 to 31 and/or the vector of claims 34 to 36 as a pharmaceutical.

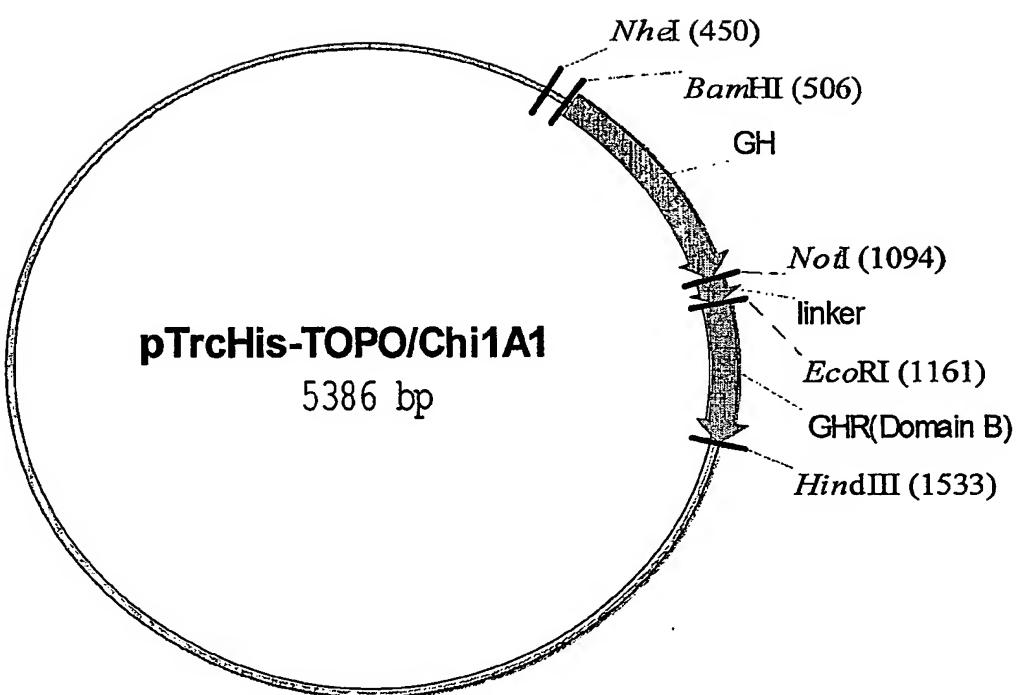
15 40. A pharmaceutical composition comprising the polypeptide according to any of claims 1 to 28, 32 or 33, and/or the nucleic acid of claims 29 to 31 and/or the vector of claims 34 to 36 wherein said pharmaceutical optionally comprises a carrier, excipient and/or a diluent.

20 41. Use of the polypeptide of any of claims 1 to 28, 32 or 33 and/or the nucleic acid of claims 29 to 31 and/or the vector of claims 34 to 36 for the manufacture of a medicament for the treatment of a disease wherein said disease is selected from the group consisting of: acromegaly; gigantism; GH deficiency; Turners syndrome; renal failure; osteoporosis; diabetes mellitus; cancer; obesity; insulin resistance; hyperlipidaemia; hypertension; anaemia; autoimmune and infectious disease; inflammatory disorders including rheumatoid arthritis.

25

42. A method of treating a human or animal subject comprising administering an effective amount of the polypeptide of claims 1 to 28, 32 or 33, and/or the nucleic acid of claims 29 to 31 and/or the vector of claims 34 to 36 and/or the pharmaceutical composition of claim 40 and/ or the medicament of claim 41 to said subject.

30

**FIGURE 1**

Name	5'—Sequence—3'
DiGHEcoF	AGGCGAATTCTTCCCAACCATTCCCTAT
DiGHNotF	CTTCAAGAGGC GGCGGCCGCTCCCAACCATTCCC TTAT
DiGHHinR	TTCCAAGCTTCATCAGAACGCCACAGCTGCCCTCCA
Lep2TrcFOR	CAAAGCTAGCGGTGGCATGCAAGT
Lep2TrcREV	AAGCTTGAATT CCTATTACGTCGACTCTAG
LepLinkFOR	CAGCTGCTGTGGCTTCGGCGGCCGCAGGTGGCGGA
LepLinkREV	AATGCCTCGAGGAATT CGGAACCTCCG
Lep2FOR	GGGAAACTCGAGGTGCCATCCAAAAAGTCCAAG AT
Lep2REV	GGGAAAGTCGACTCTAGAGCACCCAGGGCTGA GGTCC

FIGURE 2

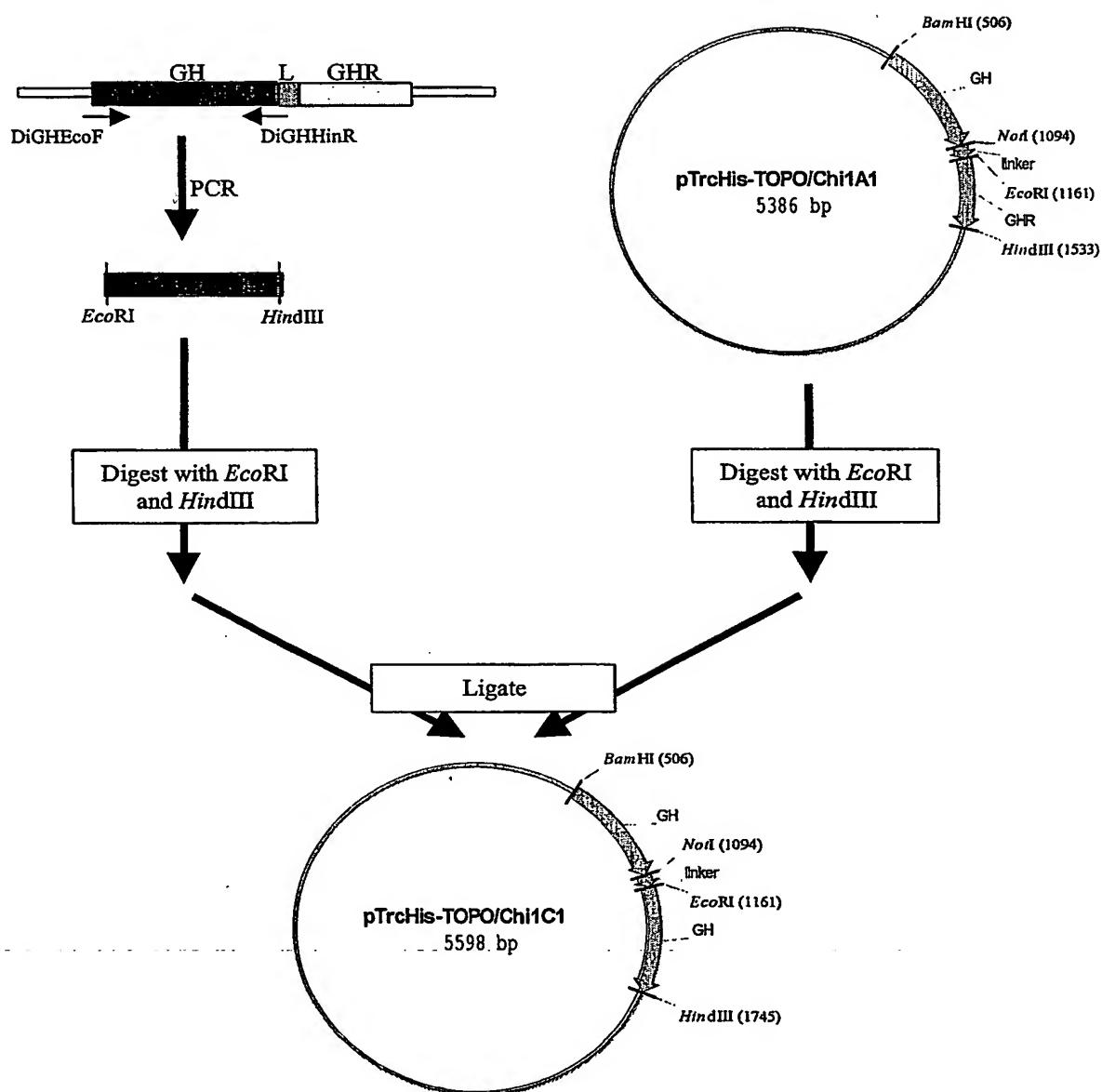


FIGURE 3

Growth hormone molecule 1

TTCCCAACCATTCCCTATCCAGGCTTTGACAACGCTAGTCTCCGC
CCATCGTCTGCACCAGCTGGCCTTGACACCTACCAGGAGTTGAAGAAG
CCTATATCCAAAGGAACAGAACAGAAGTATTCAATTCCCTGCAGAACCCCC
TCCCTCTGTTCTCAGAGTCTATTCCGACACCCCTCCAACAGGGAGGAAAC
ACAACAGAAATCCAACCTAGAGCTGCTCCGCATCTCCCTGCTGCTCATCC
AGTCGTGGCTGGAGCCCCTGCAGTTCTCAGGAGTGTCTCGCCAACAGC
CTGGTGTACGGCGCCTCTGACAGCACGTCTATGACCTCTAAAGGACCT
AGAGGAAGGCATCCAAACGCTGATGGGGAGGCTGGAAGATGGCAGCCCC
GGACTGGGCAGATCTCAAGCAGACCTACAGCAAGTTGACACAAACTCA
CACAAACGATGACGCACTACTCAAGAACTACGGGCTGCTACTGCTTCAG
GAAGGACATGGACAAGGTCGAGACATTCCCTGCGCATCGTGCAGTGCCGCT
CTGTGGAGGGCAGCTGTGGCTTC

Linker region

TTAGTGCCGCGCGCAGTCCGGCATTGGCGGCGGTGGCGGC

Growth hormone molecule 2

TTCCCAACCATTCCCTATCCAGGCTTTGACAACGCTAGTCTCCGC
CCATCGTCTGCACCAGCTGGCCTTGACACCTACCAGGAGTTGAAGAAG
CCTATATCCAAAGGAACAGAACAGAAGTATTCAATTCCCTGCAGAACCCCC
TCCCTCTGTTCTCAGAGTCTATTCCGACACCCCTCCAACAGGGAGGAAAC
ACAACAGAAATCCAACCTAGAGCTGCTCCGCATCTCCCTGCTGCTCATCC
AGTCGTGGCTGGAGCCCCTGCAGTTCTCAGGAGTGTCTCGCCAACAGC
CTGGTGTACGGCGCCTCTGACAGCACGTCTATGACCTCTAAAGGACCT
AGAGGAAGGCATCCAAACGCTGATGGGGAGGCTGGAAGATGGCAGCCCC
GGACTGGGCAGATCTCAAGCAGACCTACAGCAAGTTGACACAAACTCA
CACAAACGATGACGCACTACTCAAGAACTACGGGCTGCTACTGCTTCAG
GAAGGACATGGACAAGGTCGAGACATTCCCTGCGCATCGTGCAGTGCCGCT
CTGTGGAGGGCAGCTGTGGCTTC

FIGURE 4

Growth hormone molecule 1

FPTIPLSRLFDNASLRAHRLHQLAFDTYQEFEEAYIPKEQKYSFLQNPQT
SLCFSESIPTPSNREETQQKSNEELLRISLLIQSWSLEPVQFLRSVFANS
LVYGASDSNVYDLLKDLEEGIQLTLMGRLEDGSRTGQIFKQTYSKFDTNS
HNDDALLKNYGLLYCFRKDMDKVETFLRIVQCRSVEGSCGF

Linker region

LVPRGS PGIGGGGG

Growth hormone molecule 1

FPTIPLSRLFDNASLRAHRLHQLAFDTYQEFEEAYIPKEQKYSFLQNPQT
SLCFSESIPTPSNREETQQKSNEELLRISLLIQSWSLEPVQFLRSVFANS
LVYGASDSNVYDLLKDLEEGIQLTLMGRLEDGSRTGQIFKQTYSKFDTNS
HNDDALLKNYGLLYCFRKDMDKVETFLRIVQCRSVEGSCGF

FIGURE 5

Leptin molecule 1

GTGCCCATCCAAAAAGTCCAAGATGACACCAAAACCCTCATCAAGACAAT
TGTCAACCAGGATCAATGACATTACACACACGCAGTCAGTCTCCTCCAAAC
AGAAAAGTCACCGGTTGGACTTCATTCTGGCTCCACCCCATCCTGACC
TTATCCAAGATGGACCAGACACTGGCAGTCTACCAACAGATCCTCACCAAG
TATGCCTTCCAGAAACGTGATCCAAATATCCAACGACCTGGAGAACCTCC
GGGATCTTCTTCACGTGCTGGCCTCTCTAAGAGCTGCCACTTGCCTGG
GCCAGTGGCCTGGAGACCTTGGACAGCCTGGGGGTGTCCTGGAAGCTTC
AGGCTACTCCACAGAGGTGGTGGCCTGAGCAGGCTGCAGGGTCTCTGC
AGGACATGCTGTGGCAGCTGGACCTCAGCCCTGGGTGC

Linker region

TTAGTGC CGCGCG CAGTCCGGCATTGGCGGCGGTGGCGGC

Leptin molecule 2

GTGCCCATCCAAAAAGTCCAAGATGACACCAAAACCCTCATCAAGACAAT
TGTCAACCAGGATCAATGACATTACACACACGCAGTCAGTCTCCTCCAAAC
AGAAAAGTCACCGGTTGGACTTCATTCTGGCTCCACCCCATCCTGACC
TTATCCAAGATGGACCAGACACTGGCAGTCTACCAACAGATCCTCACCAAG
TATGCCTTCCAGAAACGTGATCCAAATATCCAACGACCTGGAGAACCTCC
GGGATCTTCTTCACGTGCTGGCCTCTCTAAGAGCTGCCACTTGCCTGG
GCCAGTGGCCTGGAGACCTTGGACAGCCTGGGGGTGTCCTGGAAGCTTC
AGGCTACTCCACAGAGGTGGTGGCCTGAGCAGGCTGCAGGGTCTCTGC
AGGACATGCTGTGGCAGCTGGACCTCAGCCCTGGGTGCTTAGTGCCG

FIGURE 6

Leptin molecule 1

VPIQKVQDDTKTLIKTIVTRINDISHTQSVSSKQKVTGLDFIPGLHPILT
LSKMDQTLAVYQQILTSMPSRNVIQISNDLENLRDLLHVLAFSKSCHL
PWASGLETLDLGGVLEASGYSTEVVALSRLQGSLQDMLWQLDLSPGC

Linker region

LVPRGS PGIGGGGG

Leptin molecule 2

VPIQKVQDDTKTLIKTIVTRINDISHTQSVSSKQKVTGLDFIPGLHPILT
LSKMDQTLAVYQQILTSMPSRNVIQISNDLENLRDLLHVLAFSKSCHL
PWASGLETLDLGGVLEASGYSTEVVALSRLQGSLQDMLWQLDLSPGC

FIGURE 7

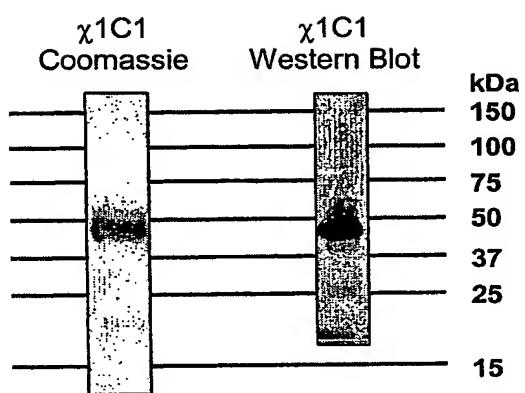


FIGURE 8

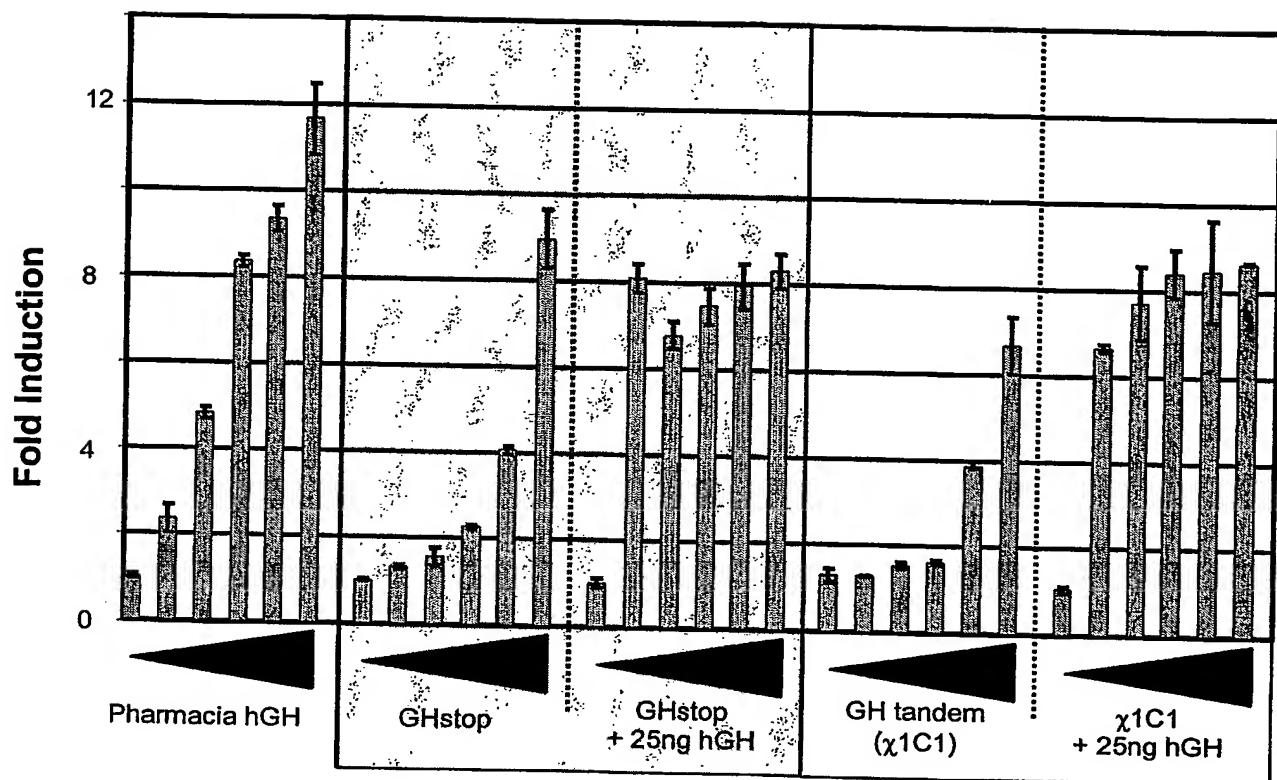


FIGURE 9

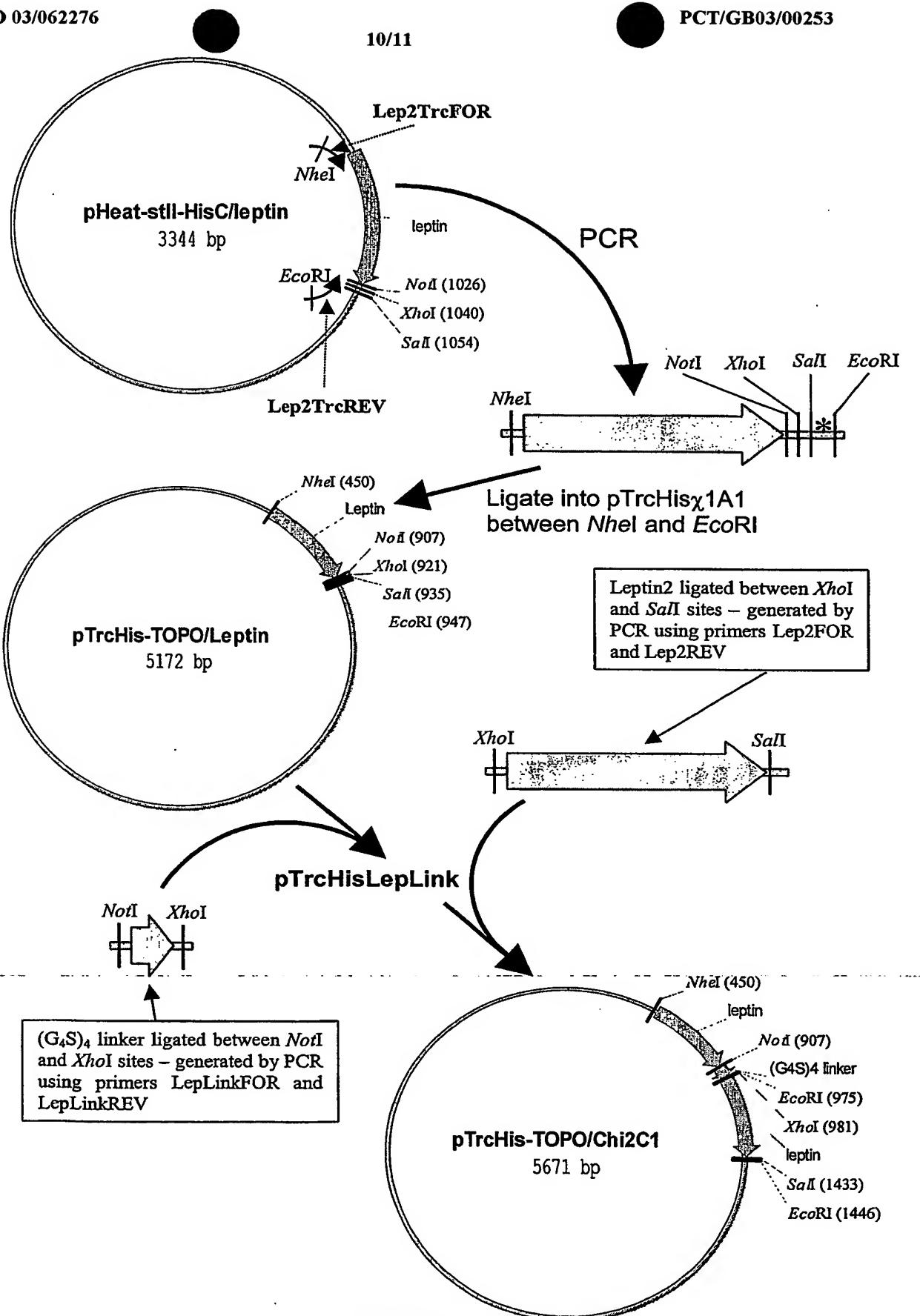


FIGURE 10

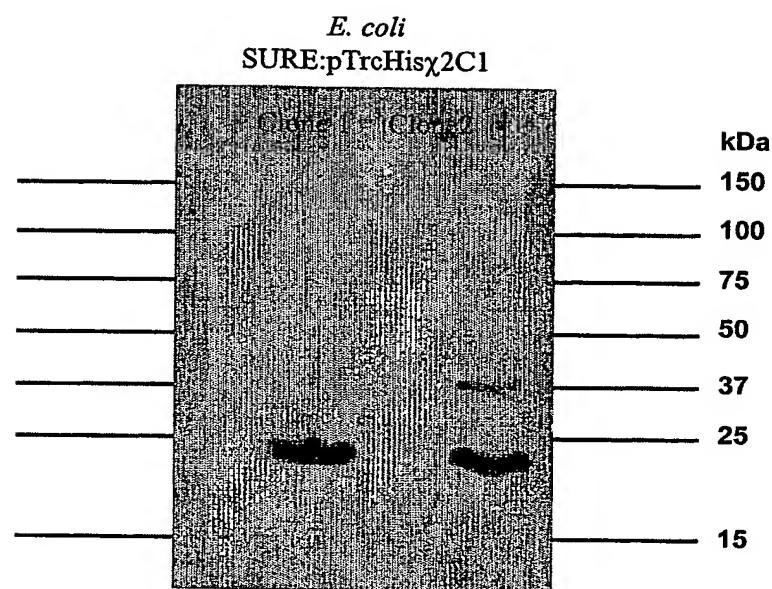


FIGURE 11

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